

STIC-Biotech/ChemLib

85106

Fr m:
S nt:
To:
Subject:

Bugaisky, Gabriele
Friday, January 24, 2003 10:19 AM
STIC-Biotech/ChemLib
09697089

RECEIVED
JAN 24 2003
STIC

amended case;
please search (including interference)

nucleic acids encoding the following regions of SEQ ID NO:2
323 C) AA 762-965 A) AA 1-88 b) AA 161-

oligomer search of n.a. encoding SEQ ID NO:2,

Also, the following, if possible

1) A nucleic acid encoding a polypeptide, wherein the polypeptide is at least 85% identical to SEQ ID NO:2, and the % identity is determined using the ALIGN program in the GCG software package, using a PAM120 weight residue table, a gap length penalty of 12 and a gap penalty of 4.

2) a nucleic acid at least 85% identical to SEQ ID NO:3, where the % identity is determined using the NBLAST program with a score of 100 and a word length of 12

thanks, gabi

Gabriele E. Bugaisky
* au 1653
* cml-10d09; mailbox 9b01
* 308-4201

Point of Contact:
Toby Port
Technical Info. Specialist
CM1 6A04
703-308-3534

Searcher: _____
Phone: _____
Location: _____
Date Picked Up: 1/29
Date Completed: 2/4
Searcher Prep/Review: _____
Clerical: _____
Online time: _____

TYPE OF SEARCH:
NA Sequences: _____
AA Sequences: _____
Structures: _____
Bibliographic: _____
Litigation: _____
Full text: _____
Patent Family: _____
Other: _____

VENDOR/COST (where applic.)
STN: _____
DIALOG: _____
Questel/Orbit: _____
DRLink: _____
Lexis/Nexis: _____
Sequence Sys.: _____
WWW/Internet: _____
Other (specify): _____